SEQUENCE LISTING

<110> Garber, Mitchell Ehren

<120> METHODS AND COMPOSITIONS FOR USE IN
    EVALUATING AND TREATING NEOPLASTIC DISEASE CONDITIONS

<130> STAN-349WO

<150> 60/629,527
<151> 2004-11-18

<150> 60/558,953
<151> 2004-04-02

<160> 13

<170> FastSEQ for Windows Version 4.0

<210> 1
<211> 20
<212> PRT
<213> human

<400> 1
Met Glu Ala Ala Asp Ala Ser Arg Ser Asn Gly Ser Ser Pro Glu Ala
 1   5   10   15
Arg Asp Ala Arg
            20

<210> 2
<211> 17
<212> PRT
<213> human

<400> 2
Glu Leu His Leu Lys Pro His Leu Glu Gly Ala Ala Phe Arg Asp His
 1   5   10   15
Gln

<210> 3
<211> 18
<212> PRT
<213> human

<400> 3
Glu Gly Glu Gly Leu Gly Gln Ser Leu Gly Asn Phe Lys Asp Asp Leu
 1   5   10   15
Leu Asn

<210> 4
<211> 17
<212> PRT
<213> human

<400> 4

Arg Glu Thr Ile Pro Ala Lys Leu Val Gln Ser Thr Leu Ser Asp Leu
1   5   10   15
Arg

<210> 5
<211> 17
<212> PRT
<213> human

<400> 5
Asp Pro Ala Lys Val Gln Ser Leu Val Asp Thr Ile Arg Glu Asp Pro
1   5   10   15
Asp

<210> 6
<211> 15
<212> PRT
<213> human

<400> 6
Arg Val Ala Ala Lys Arg Leu Lys Glu Gly Asp Thr Met Met Gly
1   5   10   15

<210> 7
<211> 19
<212> PRT
<213> human

<400> 7
Lys Thr Val Glu Ser Leu Glu Glu Thr Leu Lys Lys Ala Ser Pro Asp
1   5   10   15
Gly Tyr Asp

<210> 8
<211> 17
<212> PRT
<213> human

<400> 8
Thr Thr His Ser Ile Ser Asp Gly Lys Asp Leu Glu Lys Leu Leu Thr
1   5   10   15
Glu

<210> 9
<211> 24
<212> PRT
<213> human

<400> 9
Glu Tyr His Lys Val His Gln Met Met Arg Glu Gln Ser Ile Leu Ser
1   5   10   15
Pro Ser Pro Tyr Glu Gly Tyr Arg
            20

```
<210> 10
<211> 23
<212> PRT
<213> human

<400> 10
Arg His Gln Leu Leu Cys Phe Lys Glu Asp Cys Gln Ala Val Phe Gln
 1               5                   10                  15
Asp Leu Glu Gly Val Glu Lys
                20


<210> 11
<211> 19
<212> PRT
<213> human

<400> 11
Asp Leu Glu Val Lys Asp Trp Met Gln Lys Lys Arg Arg Gly Leu Arg
 1               5                   10                  15
Asn Ser Arg



<210> 12
<211> 30
<212> DNA
<213> human

<400> 12
ggatccatgg aagctgcaga tgcctccagg                                    30

<210> 13
<211> 71
<212> DNA
<213> human

<400> 13
accggtgtgc atcctccgcc gccgcatcct ccgccgccgg cggctggggc ttcgttggac 60
ccaatcccgt t                                                        71
```